

IDENTIFIERS

dbEST Id: 1099597
EST name: aa04a07.s1
GenBank Acc: AA455042
GenBank gi: 2177818
GDB Id: 6043155

CLONE INFO

Clone Id: IMAGE:812244 (3')
Source: IMAGE Consortium, LLNL
DNA type: cDNA

PRIMERS

Sequencing: -41m13 fwd. ET from Amersham
PolyA Tail: Unknown

SEQUENCE

GGAAGAATACAGAAATATGTTTAATACTTAGTATCAAAC TAAAAAGTAATATAA
AAAAC TTCTTTTTTTTCATGCACAGGCTTTTTCTGGTAAGGACCGCTGGGATTG
AGCTTCCGGTAAATAAGGGCCCCGTCGGCAAGACAGCATACTGCTGTCACAAGT
ACCCCTCCACCAACTGTCAATGTTGTGGTTTCTGGTATCAGTGCCAACACAGAT
AGCATGAATACTGTTGTTACCAGTGAGTTGATAATATCCAGCCGCAGCATCTTC
CCTTTCACACTGAAGCAGAAGGGGCGATGTTTTATTTTCGGCTGCACGTTATCC
TCTGCAGACCCAGCAGCAGCACTTCCCTCAACTCTTCTCAGCTGGCTGCCTGA
TCTGCGA

Quality: High quality sequence stops at base: 395

Entry Created: Jun 6 1997
Last Updated: Jun 6 1997

COMMENTS

This clone is available royalty-free through LLNL ; co
the IMAGE Consortium (info@image.llnl.gov) for further
information.

LIBRARY

dbEST lib id: 589
Lib Name: Soares_NhHMPu_S1
Organism: Homo sapiens
Organ: mixed (see below)
Tissue type: Pooled human melanocyte, fetal heart, and pregnant ute
Lab host: DH10B
Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker
R. Site 1: Not I
R. Site 2: Eco RI
Description: Equal amounts of plasmid DNA from three normalized lib
(melanocyte 2NbHM, pregnant uterus NbHPU, and fetal he
NbHH19W) were mixed, and ss circles were made in vitro
Following HAP purification, this DNA was used as trace
subtractive hybridization reaction. The driver was
PCR-amplified cDNAs from pools of 5,000 clones made fr

same 3 libraries. The pools consisted of I.M.A.G.E. cl
260232-265223, 340488-345479, and 484488-489479.

SUBMITTER

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CITATIONS

Title: WashU-Merck EST Project 1997
Authors: Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G.,
S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., M
J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Th
B., White, Y., Wylie, T., Waterston, R., Wilson, R.
Year: 1997
Status: Unpublished

MAP DATA

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IDENTIFIERS

dbEST Id: 1833454
 EST name: oz37h05.x1
 GenBank Acc: AI078580
 GenBank gi: 3412988

CLONE INFO

Clone Id: IMAGE:1677561 (3')
 Source: NCI
 DNA type: cDNA

PRIMERS

Sequencing: -40m13 fwd. ET from Amersham
 PolyA Tail: Unknown

SEQUENCE

TTTGGAAGAATACAGAAATATGTTTAATACTTAGTATCAAACATAAAAGTAATA
 TACAAAACCTTCTTTTTTTCATGCACAGGCTTTTCTGGTAAGGACCGCTGGGA
 AGAAGCTTCCGGTAAATAAGGGCCCCGTCGGCAAGACAGCATACTGCTGTCACA
 AACACCCCTCCACCAACTGTCAATGTTGTGGTTTCTGGTATCAGTGCCAACACA
 ATGAGCATGAATACTGTTGTTACCAAGTGAGTTGATAATATCCAGCCGCAGCATC
 TGGCCTTTCACACTGAAGCAGAAGGGGCGATGTTTTATTTTCGGCTGCACGTTA
 GCGTCTGCAGACCCAGCAGCAGCACTTTCCTCAACTCTTCTCAGCTGGCTGCC
 GGTTCCTGCGAAGCGATAGCAACCGCCACCGCG

Quality: High quality sequence stops at base: 410

Entry Created: Aug 10 1998
 Last Updated: Aug 10 1998

COMMENTS

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 information.

LIBRARY

dbEST lib id: 589
 Lib Name: Soares_NhHMPu_S1
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 Organ: mixed (see below)
 Tissue type: Pooled human melanocyte, fetal heart, and pregnant ute
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 same 3 libraries. The pools consisted of I.M.A.G.E. cl

260232-265223, 340488-345479, and 484488-489479.

SUBMITTER

Name:

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CITATIONS

Title:

National Cancer Institute, Cancer Genome Anatomy Project
(CGAP), Tumor Gene Index

Authors:

NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>

Year:

1997

Status:

Unpublished

MAP DATA

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